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TOPIC HIGHLIGHT

2016 Hepatocellular Carcinoma: Global view

Targeting adeno-associated virus and adenoviral gene therapy for hepatocellular carcinoma

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Abstract

Human hepatocellular carcinoma (HCC) heavily endangers human heath worldwide. HCC is one of most frequent cancers in China because patients with liver disease, such as chronic hepatitis, have the highest cancer susceptibility. Traditional therapeutic approaches have limited efficacy in advanced liver cancer, and novel strategies are urgently needed to improve the limited treatment options for HCC. This review summarizes the basic knowledge, current advances, and future challenges and prospects of adeno-associated virus (AAV) and adenoviruses as vectors for gene therapy of HCC. This paper also reviews the clinical trials of gene therapy using adenovirus vectors, immunotherapy, toxicity and immunological barriers for AAV and adenoviruses, and proposes several alternative strategies to overcome the therapeutic barriers to using AAV and adenoviruses as vectors.

Key words: Hepatocellular carcinoma; Adeno-associated virus; Adenovirus; Virus vectors

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Core tip: This review summarizes the basic knowledge, current advances, and future challenges and prospects of adeno-associated virus (AAV) and adenoviruses as vectors for gene therapy of hepatocellular carcinoma. This paper also reviews the clinical trials of gene therapy using adenovirus vectors, immunotherapy, toxicity and immunological barriers for AAV and adenoviruses, and proposes several alternative strategies to overcome the therapeutic barriers to using AAV and adenoviruses as vectors.

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INTRODUCTION

Human hepatocellular carcinoma (HCC) is the fifth leading cause of cancer death with an estimated > 33000 new cases per year in the United States, according to the American Cancer Society^[1]. In China, HCC is also one of the five most common cancers, whose incidence rate ranks fourth and mortality ranks second, with 18.43 per 10000 among all types of cancers^[2,3]. Surgery, chemotherapy, and radiotherapy are still the most common therapeutic options for HCC. Surgery is the first choice for early stage disease and offers a cure rate of 10%-20% in HCC patients. However, liver cancer recurrence rate remains high after tumor resection because of the aggressive traits of HCC, such as metastasis and chemo- or radiotherapy resistance $[4,5]$. Although tremendous efforts have been made to improve the anti-cancer effect for HCC, there are still no ideal therapeutic strategies. Gene therapy has attracted much interest as a novel promising therapeutic method since the first approved successful clinical trial for children with severe combined immunodeficiency (SCID) in 1990^[6]. Henceforth, gene therapy has rapidly developed as a strategy for single gene hereditary diseases, infections, and cancer, using various viral or nonviral vectors $^{[7,8]}$. Currently, most effective vectors are derived from recombinant viruses including adenovirus, adenoassociated virus (AAV), vaccinia virus, retrovirus, lentivirus, herpes simplex virus, Epstein-Barr virus, and chimeric viruses $[8,9]$. Since we proposed the Cancer Targeting Gene Virotherapy (CTGVT) strategy by combining cancer gene therapy and virotherapy using oncolytic adenoviral vector in 2001, numerous studies have been performed in HCC and other cancers using gene therapy^[10-12]. Additionally, due to its distinct merits, AAV is considered as the prime candidate vector for clinical gene therapy for various liver diseases $^{[13,14]}$. This review describes the advances, therapeutic mechanisms, and current challenges in using AAV and adenoviruses as vectors for gene therapy of HCC.

BIOLOGY OF AAV AND ITS APPLICATION AS A VECTOR FOR HCC

AAV, a member of the parvovirus family, has a singlestranded DNA genome of approximately 4.7 kb. The genome consists of two open reading frames (ORFs) rep and cap driven by three promoters (P5, P19 and P40), which are flanked by the 145 nucleotide long inverted terminal repeat sequences. The rep encodes four overlapping functional proteins (Rep78, Rep68, Rep52, and Rep40), which play a part in viral replication, transcriptional control, and accumulation of single-stranded progeny genomes, and the cap contains three capsid proteins (VP1, VP2, and VP3) functioning in the generation of infectious particles^[15].

Essential properties of AAV vector

Since Hermonat *et al*^[16] first used AAV as a vector for transgene delivery into cultured mammalian cells, AAV-based vectors have undergone rapid development in recent decades. Unlike the wild-type AAV that is able to integrate into the host genome, exogenous genes delivered by recombinant AAV vector can be persistently expressed in an episomal state^[17,18]. AAV vectors have some others advantages, including a broad host spectrum that allows them to infect both non-dividing and dividing cells and low pathogenicity or no cytotoxicity^[19].

Presently, there are more than 11 different serotypes of AAVs and over 100 new AAV variants to be identified and engineered into vectors. AAV2 serotype was the first to be used for the transfer of transgenes into host cells and has emerged as a promising carrier for clinical gene transfer in several single genetic disorders^[20]. As the first clinical gene therapy medicine (AAV2-LPL) authorized by European Medicines Agency (EMA ^[21], it largely promotes the development of AAV vectors for clinical transgene therapy. Besides AAV2, AAV1 and 3-9 serotypes were engineered as gene transfer vectors. AAV1 vector has a higher transfer rate in muscle than $AAV2^{[22]}$. $AAVI$, 4, and 5 can transfer genes efficiently into muscles, lung, and central nervous system $[8,23]$. The serology of AAV6 is almost identical to that of AAV1, and the AAV6 vector was designed to transfer primary human hematopoietic stem cells into a mouse model^[24]. AAV7 and AAV8 are two new members of the AAV family isolated from rhesus monkeys^[25]. Furthermore, the efficiency of gene transfer to liver with AAV7 and 8 vectors was higher than that achieved using AAV2, although a variety of host factors may influence this important parameter, such as pre-existing antibodies, gender, and transgene immunity $[25-27]$. So far, the serological profiles of AAV10 and AAV11 are not well characterized^[28,29]. AAV12 is a novel AAV serotype with transduction activity independent of sialic acid and heparan sulfate proteoglycan^[30]. The AAV variants are not further identified and exploited as vectors due to the lack of serological profiling.

AAV vector development for HCC

In recent years, many important breakthroughs have been achieved in gene therapy in several disease types, including genetic diseases and cancer. Up to January 2015, there were 127 trials of gene therapy mediated by AAV vectors registered at the Journal of Gene Medicine Clinical Trial website^[31]. These trials have been performed for the treatment of various diseases, including cancer and monogenic, neurological, ocular,

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Figure 1 Schematic of adeno-associated virus vectors packaged into adeno-associated virus serotypes 2, 8, and 9 with different promoters used for gene delivery. AFPp: Human α-fetoprotein promoter; hTERTp: A truncated human telomerase; CBA: Chicken β-actin; TBG: The liver-specific thyroxine-binding globulin; hAAT: Human α1-antitrypsin.

cardiovascular, and infectious diseases. The landmark of AAV-mediated clinical studies is the development of inherited retinal diseases and anerythrochloropsia gene therapy. AAV-mediated PRE65 gene expression efficiently recovered visual function in patients with Leber's congenital amaurosis with controlled safety and efficacy of gene transfer^[32-34]. These encouraging results widened the clinical applications of AAV vectors, including gene therapy of cancer.

AAV vectors are engineered for delivery to patients suffering from liver diseases, including familial hypercholesterolemia, viral hepatitis, and hepatic malignancies. The first gene therapy experiment for HCC using AAV vectors was carried out by Su *et al*^[35]. The constructed recombinant AAV virus, which carried the herpes simplex virus thymidine kinase (TK) gene driven by the human α -fetoprotein (AFP) enhancer and the albumin promoter, resulted in a selective killing effect on AFP-positive HCC cells but not nonhepatocyte tumor cells or AFP and albumin-negative hepatic tumor cells^[35]. Moreover, the dose required to kill the cancer cells was inversely proportional to the level of AFP expression in the cells.

Actually, besides suicide-gene-directed enzyme/ pro-drug therapy, gene therapy using AAV covers numerous therapeutic methods, such as inhibition of oncogenes and re-expression of tumor suppressor genes, immunotherapy, anti-angiogenesis therapy, and combination therapy^[19]. There are a variety of targeting strategies for AAV application, including transcriptiontargeted, receptor-targeted, and conjugate-targeted strategies and various alternative AAV serotypes^[8]. The application of human telomerase reverse transcriptase (hTERT) promoter was a good candidate for transcription-targeted cancer gene therapy^[13]. In general, cancer cells have high telomerase activity, and hTERT, a catalytic subunit of the telomerase, is transcriptionally upregulated exclusively in about 90%

of cancer cells^[36]. Thus, telomerase is an excellent tumor marker. We first constructed a novel AAV vector targeting telomerase activity and investigated its targeting capability and anti-tumor potential though carrying a human interferon- β gene^[13]. The recombinant virus AAV-hTERT-human interferon (hIFN)-β displayed cancer-specific hIFNβ expression and cytotoxicity in various human cancer cells *in vitro* and suppressed tumor growth in nude mice of lung cancer A549 and colorectal cancer SW620 xenografts^[7]. In addition, we and others generated a recombinant AAV vector containing the tumor necrosis factor alpha related apoptosis inducing ligand (TRAIL) gene under the control of the hTERT promoter. The AAV-hTERT-TRAIL virus exhibited cancer-specific cytotoxicity and apoptosis that which significantly suppressed the growth of HCC xenograft tumors^[37,38]. These results indicated that AAVs in combination with hTERT-mediated therapeutic gene expression provide a promising targeting approach for developing effective therapy for HCC (Figure 1).

AAV-mediated gene therapy of HCC has made great progress in other research areas. Intraportal injection of recombinant AAV carrying kallistatin gene suppressed hepatic and subcutaneous HCC tumors through antiangiogenic and antiproliferative activities^[39]. Ling *et al*^[40,41] validated that AAV serotype 3 is an excellent vector in efficiently transducing human liver cancer because AAV3 uses human hepatocyte growth factor receptor as a cellular co-receptor for binding and entry into these cells, which implies that AAV3 vectors can be applied to gene therapy of liver cancer. In addition, AAV8 may be the best liver-specific transfer vector and has good prospects for liver cancer gene therapy^[42]. In particular, RNA interference (RNAi)-based approaches, such as antisense hypoxiainducible factor-1 α and microRNA (miRNA)-targeted therapy mediated by AAV have been recently ex-

ploited as new anti-cancer treatments for HCC^[43-45]. Systemic administration of AAV-mediated miR-26a delivery efficiently inhibited HCC cell proliferation, induced tumor-specific apoptosis, and suppressed tumorigenesis in a murine liver cancer model^[45]. Other than the general strategy for miRNA replacement therapies, inhibiting the oncogenic miR-221 by miRNA sponge was developed for therapy of HCC, in which AAVs were genetically modified to drive the expression of multiple binding sites for miR-221 $^{[44]}$.

Combination therapy is an important tactic for clinical cancer therapy. AAV-mediated gene therapy is widely considered as a potential adjuvant of other therapies. We attempted combination therapy with AAV-hTERT-TRAIL and cisplatin, which could have a synergistic therapeutic effect on HCC. As expected, treatment with both AAV-hTERT-TRAIL and cisplatin exhibited a stronger inhibitory effect and induced more significant apoptosis compared with either agent alone in HCC cells and animal tumors^[46]. Other studies showed that radiotherapy can enhance transduction of HCC cells by recombinant AAV *in vitro* and *in vivo*[47]. Even the cocktail viral gene therapy, combining the effect of AAV transducing hepatocyte growth factor dringle 1 and adenovirus transducing p53, significantly induced tumor cell death, inhibited tumor angiogenesis and tumor growth, and prevented tumor metastasis in HCC models^[48]. Multigene-based combination therapy is an effective practice in cancer gene therapy. AAVmediated coexpression of apoptin and interleukin (IL)-24 in HCC significantly suppressed the growth and induced apoptosis of HepG2 cells *in vitro* and in xenograft nude mice^[49]. An AAV6 serotype designed for dendritic-cell-based cancer immunotherapy can be a useful targeting approach for efficient HCC therapy^[50], which could lay the foundation for further development for AAV-mediated HCC immunotherapy.

THERAPEUTIC APPLICATION OF ADENOVIRUS IN HCC

Adenovirus has been the most common gene transfer vector for cancer gene therapy in past decades because of its unique advantages, such as broad tropism for infecting many human tissues including hepatocytes, capability of transducing nonreplicating cells and replicating cells, and easy acquisition of high titers, benefiting clinical use and efficient transgene $expression^{[51,52]}$. Numerous studies have reported the potential application of adenovirus-mediated gene therapy for a wide variety of diseases and indicated their beneficial effects, tolerability, and safety. The following three aspects describe oncolytic adenovirus vector, adenovirus-mediated immune treatment, and clinical trials for HCC therapy, respectively.

Oncolytic adenovirus vector

Currently, oncolytic viruses represent a group of pro-

mising anti-cancer agents with the ability to lyse infected cancer cells but not normal cells. The oncolytic viruses include genetically modified adenovirus, vaccinia virus, herpes simplex virus, and reovirus $[10]$. ONYX-015, an oncolytic adenovirus designed with the E1B 55-kDa gene deletion, was engineered to replicate in and lyse cancer cells selectively^[53]. Clinical trials using ONYX-015 alone or in combination with chemotherapy have been widely performed in patients with head and neck cancer, which achieved an obvious anti-cancer effect^[54,55]. In addition, many targeted strategies based on oncolytic adenoviruses were conceived, and they exhibited potent anti-tumor activity in various preclinical studies $^{[11,56,57]}$. Previously, our CTGVT, which combined the superiority of gene therapy and virotherapy, brought new hope for cancer therapy^[10,58]. Similar to ONYX-015, the novel oncolytic adenovirus ZD55 was first engineered with E1B55-kD protein deletion based on CTGVT, and was combined with anticancer gene therapy to form the ZD55-gene system^[58]. Our many experiments confirmed that the ZD55 gene exerted a potent anti-tumor effect in multiple tumor cell lines and mouse models through the synergetic mechanisms of the oncolytic action of the virus itself and overexpression of anti-tumor genes^[11,52,59-63]. In particular, the novel oncolytic adenovirus ZD55- Smac increases anti-tumor activity of ZD55-TRAIL against HCC *via* a synergetic apoptotic effect^[64]. In general, HCC frequently displays a high resistance to TRAIL-mediated cell death due to high expression of inhibitor of apoptosis proteins (IAPs), while Smac strongly inhibits IAPs and increases sensitivity of HCC cells to TRAIL^[65]. Thus, combination treatment with ZD55-Smac and ZD55-TRAIL led to rapid and potent activation of apoptosis in HCC cells and eliminated completely tumor xenograft in all treated animals, which could provide a useful strategy for therapy of $HCC^{[64]}$.

Furthermore, numerous other combination strategies were adopted in HCC therapy based on the oncolytic adenovirus ZD55 system. Chu et al^[62] found that adenoviral vector expressing cylindromatosis (CYLD) augments anti-tumor activity of ZD55-TRAIL by suppression of nuclear factor-κB survival signaling in HCC. Pan *et al*[66] constructed ZD55-mediated X-linked inhibitor of apoptosis protein (XIAP)-shRNA by RNAi to inhibit high XIAP of HCC cells and concluded that combination of ZD55-shXIAP and ZD55-TRAIL led to synergistic anti-tumor activity in experimental HCC. Moreover, the combination of XAF1, IL-24, or Smac-armed ZD55 and chemotherapy exhibited significantly enhanced suppression of HCC growth by a synergistic mechanism, especially in the induction of apoptosis^[67-69].

However, traditional oncolytic adenovirus lacks the ability to target liver cancer. To improve the anti-cancer effect of oncolytic adenovirus in liver cancer, three main strategies have been developed in recent years. First, the transcription-targeted strategy is designed

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CBA: Chicken β-actin; TBG: The liver-specific thyroxine-binding globulin; hAAT: Human α1-antitrypsin.

using cancer- or tissue-specific promoter to control the expression of viral genes essential for replication^[70,71], which results in selective expression of viral genes, propagation of virus progeny, and tumoricidal activity^[9,10]. We designed AFP-regulated oncolytic adenovirus AD and obtained a better anti-cancer effect in HCC than other types of cancer^[72-76]. Golgi protein (GOLPH2) is an excellent HCC marker $[77,78]$, therefore, we constructed a novel GOLPH2-regulated oncolytic adenovirus GD55 that targets $HCC^{[79]}$. We showed that the novel GD55 had higher adenovirus replication ability and infectivity in liver cancer cells than the common oncolytic adenovirus ZD55. In addition, GD55 exerted a significant growth-suppressing effect on HCC cells or xenograft but caused little damage to normal liver cells, which may provide a promising oncolytic virus for future liver cancer treatment^[79]. The second strategy targets the tumor signaling pathway through deletion or mutation of key adenovirus genes or some bases that are necessary for adenovirus replication in normal cells but not in tumor cells. The classical design of oncolytic adenovirus is ONYX-015 and ZD55, whose E1B 55 kDa gene is deleted by genetic engineering^[53,58]. They are supposed to target and lyse p53-dysfunctinal tumor cells preferentially but not adjacent normal cells^[80], but further study demonstrated that the adenovirus mutant enhanced the viral mRNA late nuclear transport and oncolysis^[81]. We also noted inhibition of liposarcoma by histone deacetylase inhibitor occurs irrespective of p53 mutational status but *via* targeting of the MDM2-p53 signaling axis and phosphatase and tensin homolog^[82]. Another modification was a 24 base pair deletion in the E1A region, such as oncolytic virus Ad5-E1A(Δ24); which is responsible for binding retinoblastoma (Rb) protein, and its replication is restricted in Rb-inactive arrested cells and exhibits tumor-selective capability^[56,71,83]. The third approach is the receptor-targeted or capsidmodified strategy. Adenovirus can efficiently infect host cells by binding to specific receptors on the target cell surface with fibers on the capsid. Thus, modification of the adenovirus capsid may improve the binding ability of adenovirus to target cells. The adenovirus vector with genetically modified fibers (RGD-4C or chimera fibers of different serum types) demonstrated expanded tropism *via* utilization of a coxsackievirus and adenovirus receptor-independent cell entry mechanism^[84,85]. Otherwise, efficient and selective gene transfer into primary human tumors using single-chain antibody-targeted adenoviral vectors with native tropism abolished the specific targeting ability^[86]. In Table 1, we sum up Liver- or hepatocellular carcinoma- specific promoters and delivery gene in adeno-associated virus vectors for the past few years.

Adenovirus-mediated immunotherapy

The three traditional anti-cancer therapies (surgery, radiation, and chemotherapy) often carry risks and/or cause adverse side effects and show limited efficacy, particularly for late-stage cancer. The fourth option is cancer biotherapy, including oncolytic viruses and immunotherapy, which has emerged as a promising therapy in preclinical trials and cancer patients. Immunotherapy was considered the Breakthrough of the Year for 2013 and 2014 because of the efficacy of antibodies against cytotoxic T-lymphocyte-associated protein (CTLA)-4, programmed death (PD)-1, ligand 1, and chimeric antigen receptor^[96,97]. Encouragingly, CD19-targeted T cells achieved complete remission in children and adults with chemotherapy-refractory acute lymphoblastic leukemia^[98]. These also brought inspiration and confidence to acquire the expected therapy effect either *via* oncolysis or antitumor immunity by recombinant oncolytic adenovirus.

Fortunately, people began to realize that oncolytic adenoviruses cause the immune system to stimulate an anti-tumor immune response^[99]. A novel oncolytic adenovirus, Ad5D24-CpG, was engineered by inserting 18 immunostimulatory islands into Ad5D24. This virus showed increased anti-tumor activity *via* the stimulation of Toll-like receptor 9 and inactivation of myeloid-derived suppressor cells in modified virustreated mice^[100]. Moreover, to achieve superior anticancer immunity, oncolytic adenoviruses are often designed to express immunostimulatory molecules including CD40L, IL-2, IL-12, IL-24, and granulocytemacrophage colony-stimulating factor (GM-CSF) $^{[101]}$. Among these constructs, oncolytic adenovirus coding for GM-CSF (Ad5-D24-GMCSF) was a typical agent, and it induced anti-tumor immunity in cancer patients with advanced solid tumors refractory to standard therapies, indicating that the treatment was safe. The

TACE: Transcatheter arterial chemoembolization; ADV-TK: Adenovirus thymidine kinase; AdVhAFP: α-fetoprotein adenoviral vector; GM-CSF: Granulocyte-macrophage colony-stimulating factor; TK99UN: Adenoviral vector containing thymidine kinase of herpes simplex virus; im: Intramuscularly; it: Intratumoral; id: Intradermally.

tumor completely disappeared in $2/20$ patients^[102].

Further modifications were made in the following approaches to improve clinical anti-tumor immunological benefit. The adenovirus serotype 5 capsid was replaced with serotype 3 fiber knob to form chimeric adenovirus vector Ad5/3-D24-GMCSF, avoiding the problem of coxsackie-adenovirus receptor downregulation in advanced tumors^[99]. Another immunostimulatory molecule of interest is CD40L, a multifunctional protein. Oncolytic adenovirus encoding CD40L led to a strong anti-tumor effect $[103]$. Efficient targeted cancer immunotherapy was achieved with oncolytic adenovirus coding for a fully human monoclonal antibody specific for CTLA-4 (Ad5/3-Δ24aCTLA4), avoiding the severe immune-related adverse events by systemic administration of monoclonal antibodies ipilimumab or tremelimumab blocking CTLA- $4^{[104]}$. It also suggests the feasibility of immunotherapy with oncolytic adenovirusmediated CTLA4 antibody^[105].

An alternative means of anti-tumor immunotherapy is oncolytic adenovirus-vector vaccines. Orally delivered oncolytic adenovirus vaccines have been utilized to prevent adenovirus-induced respiratory illness in military recruits, demonstrating safety and high efficacy^[106]. The experience suggested that oral administration of live oncolytic adenoviruses holds promise for immunization against liver cancer and other infectious diseases because live adenoviruses can express intact tumor-associated antigens as transgenes in infected cells^[107]. Another novel approach is incorporation of antigenic epitopes into adenovirus capsids, eliciting the strongest humoral and cellmediated immune responses, both prior to and during virus replication, against cancer and infection^[106,108,109].

Clinical trials for adenovirus-associated HCC therapy

The first gene transfer with recombinant replicationdefective adenovirus was successfully implemented in HCC cells in 1995, which led to induction of sensitivity to ganciclovir in human HCC cells by adenovirusmediated herpes simplex virus TK gene $[110]$. Since then, 293 papers have been published in the field of adenoviruses and HCC, and 317 gene therapy studies using adenovirus vectors have been registered at ClinicalTrials.gov. Adenoviruses are the most frequently used gene transfer vectors in clinical trials according to the Journal of Gene Medicine Clinical Trial site^[31]. However, at the time of the preparation of this review (May 2015), official sources listed only eight clinical trials that described the status (Table 2), efficacy, and safety of adenovirus-associated therapy in HCC.

In particular, there are two studies that are currently recruiting participants. Recombinant human adenovirus type 5, with an E1B gene deletion, combining transcatheter arterial chemoembolization (TACE), is being tested as a stand-alone therapeutic intervention in a phase Ⅲ trial in patients with advanced HCC not amenable to surgery or local ablative therapy (NCT01869088). A phase Ⅱ trial of rAd-p53 artery injection combined with TACE in adults with HCC is being sponsored by Shenzhen Sibiono Genetech (NCT02418988). rAd-p53 was the first approved adenovirus agent worldwide for the treatment of head and neck cancer, and great success was achieved in these patients, especially when combined with chemotherapy or radiotherapy. In addition, three clinical trials have recently been completed using adenovirus vectors. The preliminary safety and efficacy of double-dose adenovirus-mediated adjuvant therapy (Adv-TK, adenovirus expressing TK) was evaluated in phase Ⅱ trials, resulting in improved outcome of liver transplantation in patients with advanced HCC (NCT02202564). In addition, liver transplantation with Adv-TK gene therapy improved survival in patients with advanced HCC (NCT00300521). Intratumoral injection of TK99UN (an adenoviral vector containing TK) was assessed in a phase I clinical trial in adult HCC patients (NCT00844623). Two trials using adenovirus vectors were terminated for undisclosed reasons. A phase I trial of the effectiveness of gene therapy with Ad5CMV-p53 was also terminated in patients with liver cancer that could not be surgically removed (NCT00003147). A suspended phaseⅠ/Ⅱ trial is testing immunization, safety, and toxicity of AFP plus GM-CSF plasmid prime and AFP-armed adenoviral vector (Adv-hAFP) in patients with locoregionally pretreated HCC (NCT00669136). A vaccine therapy study using Adv-hAFP was halted prior to enrollment

for treating patients with stage Ⅱ, ⅢA, ⅢB, or ⅣA liver cancer (NCT00093548).

EXISTING PROBLEMS AND CHALLENGE

Toxicity and immunological barriers for AAV

The nonpathogenic feature of AAV endows it as a promising gene therapy vector with little or no acute toxicity to the host. Results from gene therapy trials with AAV vectors, especially some exciting results from clinical studies of hemophilia B, congenital blindness, and familial lipoprotein lipase deficiency, have confirmed their therapeutic potential^[110,111].

However, some of the limitations of *in vivo* AAV gene transfer have emerged. First, the host immune response to AAV capsid is an important obstacle to safety and efficacy of AAV-vector-mediated gene transfer *in vivo*[112,113]. AAV2-capsid-specific cytotoxic T cells were detectable following AAV2-mediated hepatic delivery of factor IX in hemophilia B patients, which resulted in killing and clearance of transduced hepatocytes and affected the therapeutic efficacy. It was hypothesized that this was caused by rejection of transduced hepatocytes by AAV-capsid-specific memory CD8⁺ T cells reactivated by AAV vectors, because these patients harbor a population of capsidspecific memory cytotoxic T cells formed during childhood infection with wild-type AAV2^[114]. Second, the humoral immune response is a universal obstacle in virus-mediated gene transfer *in vivo*, largely affecting the therapeutic efficacy. Conceivably, both the transgene protein and AAV capsid can produce relevant antibodies. Anti-AAV capsid neutralizing antibodies are highly prevalent and detectable in two-thirds of the population. Even high titer AAV neutralizing antibodies can completely inhibit vector transduction to the target tissue, leading to lack of efficacy^[113]. Although the integration potential of AAV into the host genome offers long-term transgene expression in animal experiments or clinical trials, there is a risk of insertional mutagenesis that may induce carcinogenesis^[115]. In particular, a high incidence of HCC was observed in mice or other mouse models after systemic delivery of AAV gene therapy vector $^{[115]}$. The hepatic genotoxicity may be caused by AAV integration into the RNA imprinted and accumulated in the nucleus (Rian) locus, resulting in overexpression of proximal miRNAs and retrotransposon-like 1 associated with $HCC^{[15]}$.

Problematic limitations for oncolytic adenovirus

Adenovirus has a broad range of vertebrate hosts, including humans, and commonly causes illnesses such as mild respiratory infections and cold-like syndrome in young children. Although wild-type adenovirus can kill some cancer cells, it also has many side effects. Numerous modified oncolytic viral constructs, such as H101, Ad-p53, and Ad5-D24-GMCSF, have indicated potent anti-tumor efficacy in patients with solid cancers refractory to standard therapeutics with limited or no toxicity to normal tissue^[9]. However, there currently are some inevitable obstacles for clinical application of systemic adenovirus-mediated gene therapy. These are high prevalence of neutralizing antibodies, induction of immune and inflammatory responses, high promiscuity due to widespread expression of the coxsackie-adenovirus receptor, and adenovirus sequestration by the liver $[116]$. The approximately 36 kb genome of adenovirus comprises a variety of structural, replication, and regulatory genes, resulting in the complexity and uncertainty of the toxic effect induced by oncolytic adenovirus during systemic administration. The representative oncolytic adenovirus mutant is ONYX-015 (designed by Onyx Pharmaceuticals). Although phase I and II clinical trials of ONYX-015 were completed in patients with various solid tumors, a phase Ⅲ trial for the treatment of recurrent head and neck cancer patients was suddenly repealed because of possible safety problems a decade ago. Therefore, improving anticancer efficacy and reducing toxic effects and immune response to adenovirus vectors remain potential challenges to successful HCC therapy.

FUTURE PROSPECTS

With the rapid development of HCC incidence and mortality in China, there is an urgent need for innovative, alternative therapies for HCC patients. Despite anti-tumor efficacy being achieved by AAVor adenovirus-mediated gene therapy in experimental liver cancer models, researchers soon realized its limitations and ongoing efforts are being made to resolve these limitations. Efficient transfer of genes/small RNAs to the majority of cancer cells is still unrealistic for solid tumors $[117]$. To date, there are 17 trials using AAV vectors and 154 trials using adenoviruses for gene therapy of cancer registered at the Wiley Clinical Trial site $[31]$, although none of these trials for AAV and only eight for adenoviruses are investigating liver cancer. Therefore, more efficient AAV or adenovirus vectors targeting HCC should be designed to achieve successful treatment of liver cancer. Current strategies are mainly aimed at chemical modification of the virus capsid, serotype substitution of different virus types, and hybrid vectors combining viral and synthetic vectors to improve therapeutic efficacy for HCC. These new strategies have gradually demonstrated that the modified vectors have the ability to escape neutralizing antivirus antibodies, to overcome liver tropism, and to reduce humoral and cellular immune responses and liver toxicity even after systemic virus administration, while maintaining their natural biological activity $[118]$. In addition, the various combination strategies between different virus vectors or gene therapy and conventional/cell therapy can optimize the efficacy of AAV or adenovirus-mediated therapy. Thus, we believe that optimal scheduled

combinatorial regimens will likely have promising antineoplastic effects in the field of gene therapy with modified virus vectors.

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